| Fri May 28 09:30:31 2004 | align505_20020028508 | Page 1 |
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| GenCore version 5.1.6 Copyright (c) 1993 - 2004 Compugen Ltd. | 333 355 76554 | US-09-790-264-29 Sequence 29, Appl US-09-790-264-41 Sequence 41, Appl US-09-790-264-43 Sequence 43, Appl US-09-790-264-31 Sequence 31, Appl |
| OM nucleic - nucleic search, using sw model | 38 24.8 1.4 1440 39 24.4 1.4 1350 | Sequence 40, Sequence 35, |
| Run on; May 19, 2004, 15:48:38 ; Search time 25 Seconds (without alignments) 4.070 Million cell upda | C 40 24 1.4 846 C 41 22.6 1.3 825 C 42 22.6 1.3 189 C 43 22.6 1.3 384 | Sequence 13, Sequence 14, Sequence 45, |
| | C 44 22.4 1.3 1384 C 45 21.4 1.2 1338 | Sequence 37, Sequence 3, |
| | aat 1738 | ATTOMARNING |
| Scoring table: IDENTITY_NUC Gapop 10.0 , Gapext 0.5 | - | |
| Searched: 42 seqs, 29275 residues | RESULT 1 US-09-790-264-9 | |
| Total number of hits satisfying chosen parameters: 84 | ; Sequence 9, Application US/09/90264 ; Patent No. US200208508A1 | 190264 |
| Minimum DB seq length: 0 Maximum DB seq length: 2000000000 | APPLICANT: HOLEANAI, Douglas APPLICANT: Goodearl, Andrew | A. D.J. |
| Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries | TITLE OF INVENTION: NOVEL GENES E TITLE OF INVENTION: PROGNASTIC, J TITLE OF INVENTION: PROGNASTIC, J | NES ENCODING PROTEINS HAVING TIC, DIAGNOSTIC, PREVENTIVE, THERAPEUTIC, AND OTHER |
| Database : US20020028508A1.seq:* | CURRENT APPLICATION NUMBER: US/09/790,2 | US/09/790, 264 |
| Pred. No. is the number of results predicted by chance to have score greater than or equal to the score of the result being p and is derived by analysis of the total score distribution. | a ; PRIOR AF rinted, ; PRIOR FI ; PRIOR AF | 09/065,661 3 09/298,531 |
| \$ SUMMARIES | | 309/065,363 |
| Result Query No. Score Match Length DB ID Descrip | PRIOR PRIOR | 209/337,930 200/100 705 |
| 85.4 1512 1 US-09-790-264-9 77.7 1475 1 US-09-790-264-16 77.7 1475 1 US-09-790-264-22 | 9, Appli 9, PRIOR FILING DATE 16, Appli 9, PRIOR FILING DATE 22, Appl 9, PRIOR FILING DATE | 99/363,630 |
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| 31.6 1.8 1353 1 03-09-790-264-19 31.6 1.8 1475 1 03-09-790-264-16 31.6 1.8 1475 1 03-09-790-264-20 31.6 1.8 1475 1 03-09-790-264-20 31.6 1.8 1475 1 03-09-790-264-20 | 19, Appl Qy 143 16, Appl Qy 143 16, Appl Db 18 | CTCAGGAGGTGCCTCCAGGCGGCCGAGTGGGCCCTGAGGCCCAGCAAGGGCTAGGGTCCAT 202 |
| 25 30.8 1.8 1311 1 US-09-790-264-31 26 30.8 1.8 1440 1 US-09-790-264-40 27 30 1.7 140 1 US-09-790-264-40 28 30 1.7 30 1 US-09-790-264-25 28 30 1.7 1350 1 US-09-790-264-32 | 31, Appl Qy 203 40, Appl Db 78 | CTCCAGTCCCAGGACACAGCAGCGACGACGACGACGACCAGCGCTGGGCTCCAGCAGCATCAG 262 |
| 29 29.4 1.7 1384 1 US-09-790-264-37 30 29.4 1.7 1452 1 US-09-790-264-41 31 29.4 1.7 2148 1 US-09-790-264-33 | 37, Appl Qy 263 41, Appl Db 138 | CAGCCCCCAGAGACCGGGGAGGACAGGTCGCCCCACCACCACCGGAGGAGCAGCTCCTGCC 322 |
| 32 27.6 1.6 249 1 US-09-790-264-18 33 27.6 1.6 630 1 US-09-790-264-24 | 18, Appl 24, Appl Qy 323 | CCTGTCCGGGGGATGACTGATTCTCCTCCGCCAGGCCACCCAGAGAGAAGGCCACCCCG 382 |

THERAPEUTIC, AND OTHER н С 1522 1397 1457 144 105 165 264 1278 CTGGGATCTTCTCTGTGAATCCACCCCTGGCTACCCCCACCGCTACCGCGAT 1337 204 eccoccaseascesses and a second concerned a second and a second and a second a se 106 CAGGAGGTGCCTCCAGGGGGGCCAGTGGGGCCTGAGGGCCCAGGGTAGGGTCCATCT 205 CCAGTCCCAGGACACCAGCGGCCACCATGGCCACGCTGGGGCAGCAGCAGCAGCA 145 CAGGAGGTGCCTCCAGGCGGCCAGTGGGGCCTGAGGGCCCAGCAAGGGCTAGGGTCCATCT CCCAAGGCCAGGTGGGCCCTCAGCTGAGGGAAGGTACGAGCTCCCTGCAGGCCTGGAG 1338 cccaageccagereecccrcaecreagecaagecraceracereese Gaps Indels 122; Score 1350; DB 1; Length 1475; Pred. No. 7e-123; 0; Mismatches 0; Indels 122 built of the set Query Match 77.7%; Best Local Similarity 92.1%; Matches 1421; Conservative ; FEATURE: ; NAME/KEY: CDS ; LOCATION: (194)...(442) US-09-790-264-16 TYPE: DNA ORGANISM: Homo sapiens 265 1463 RESULT 2 ዋ 8 g 8 δ g 8 B å 8 R à 9 g CTGGGATCTTCTCTGTGATCCCCCGGGCTACCCCCCGGGCTACGGCCAT 1462 1222 1282 1157 1218 critchechertertechecertechessassertecessassertessastes 1277 983 AATGAAGGAAGAAGTGCAGAGGCTGCAGGTCCAGGGTGGACCTGCTGGAGGAGGAGGAGCTGCA 1042 1102 1162 978 coccescasecerecreatecatericertecatecatecatececerecatecatecereatecea 1037 1097 1283 CCAACATGCTGGGGGTCCAGGAAGCCACCTCGGGGGTGACTGAGCGGGAAGGCCAGGGGCC 1342 1343 CTTCCTCCTCCTCCTCCCCTTCCTCGGGGAGGCTCCCCCGGACGCCATGGGATGGG 1402 977 677 857 917 742 502 562 622 497 682 557 617 802 737 863 COCCOCCAGTACTGGTGCCAGTGTTGGGAGGGGGCACAGCCTGTCTGCAGGGCGGTACACT 922 cecceecacitracreereccaciterreceacececaciectercrecterecteracecacit 797 CTGTGTGCCCAAGGGAGGGCCCCCAGGGTGGCCCCCAACCCGACAGGAGTGGACAGTGC 982 377 378 reacededecercerererererectederectedecercerererectedecererereded 437 442 317 1038 GCAGATTTCCTTCCTGGAGGAGCAGCTGGGGGGCCCTGCTGCTGCTAGAAAGACTCGTGACT 1098 GCC2AGGGCCCAGGCTGAACTGAGGCCCCTCACGCGCGCCCTGCAGCCCCTGCCCTGC 798 crierereccaaeeecccccccaeeereecccccaaeereeccccaaccceaereeacaeree 043 GCTGGTGCTGGCCCCACTGCCTGGCCTCGCAGGCACTGGAGCATGGGCTCCCGGA écréerécrescentecacresceresceresceresceresceres de la constance de CCCCGGCAGCCTCCTGGTGCACTCCTAGCAGCTCGGCCGCATCGACTCCCTGAGCGA GCAGATTTCCTTCCTGCAGCAGCAGCTGCGGGGTCCTGCTGCTAGCAAGAAGACTCGTGACT 558 CAGCGGGGCTTCCTGGGGCCTGTGGAGCAGCAATATGCCAGCCGCCCATGCCGGAACGGAGG 858 AATGAAGGAAGAAGTGCAGAGGCTGCAGGTCCAGGGTGGACCTGCTGGAGGAGGAGGAGCTGCA GGCAGTGGGCGCACAGAGCACGCCTACCGGCCCGGGCCGTAGGGTGTGTGCTGTCCGGGC 438 creccadesecadesecrecadecadecadecadecadecadecadeseccedesec CAGCCCTGGGCTGGCCTGGCCTGGCTTGGCTGCTGCTGGCTGGAAGGAGC CAGCGGGCTTCCTGGGGCCTGTGGAGCAGCAATATGCCAGCCGCCATGCCGGAACGGAGG 743 GAGCTGTGTCCAGCCTGGCCGCTGCCGCTGCCCTGCAGGATGGCGGGGTGACACTTGCCA 618 cadecrearecreacecreccecreccecreccreccaecarecreaterecte CCTGGAGGCACCACCATGAGGGGCCTCTCAGGAGGTGCTGCTGATGTGGCTTCTGGTGTT 258 CCTGGAGGCACAGGCCATGAGGCCCCCCAGGAGGTGCTCCTGGTGTT 503 TCACGGGGACCCTGTCCGAGTCGTTCGTGCAGCGTGTGTACCAGCCCTTCCTCACCAC 563 CTGCGACGGGCACCGGGCCTGCAGCACCTACCGAACCATCTATAGGACCGGCCTACCGCCG 1403 738 1163 623 683 803 923 918 1103 1223 383 443 **1**98 9 3 a 8 g 8 a 3 g 8 ß à g 8 8 8 đ \mathcal{S} g 8 B: 3 ß 8 8 ਨੋ 8 ą ð 8 B à

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publication No. US20030166907A1
GENERAL INFORMATION: Query Match 99.9%; Best Local Similarity 99.6%; Matches 272; Conservative : Query Match 99.9%; Best Local Similarity 99.6%; Matches 272; Conservative TYPE: PRT ORGANISM: Homo sapiens US-09-852-472-2 DUITIA -09-852-472-2 LENGTH: 273 121 181 RESULT 3.04 PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR PRIOR -sn $\hat{\sigma}$ a \mathcal{S} q 8 q đ \mathfrak{F} å 8 đ 8 A 6 \mathcal{S} g

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| | | UDSLASU. | TI III | NETION: ATION: | , Avi | Botstein, David | cs, Luc Dan | Ferrara, Napoleon Filvaroff, Ellen | Fong, Sherman Gao Wei-Diang | Hanspet | en, mary , Audrey | L, Paul . L. J. Ch | Gurney, Austin L | , Ivar J | Kuo, Sophia S. Napier, Marv A. | James; | NICNOLAS F argaret Ann | Shelton, David L. Stewart, Timothy A. | Daniel a P Mi | Wood, William I | Acids | 30P1C18 NUMBER: | : 2002- | UMBER: U 2001-07- | UMBER: 6 | UMBER: 6 | UMBER: 6 | 1997-11- ITMBER: 6 | 1997-11-21 | 1998-03-10 | NUMBER: 60 | NUMBER: 6 | UMBER: 6 | 1998-03- TIMBER: 6 | 1998-03- | | UMBER: 6 1998-03- | TUMBER: 60/078936 1998-03-20 |
| SPRVAPNE SPRVAPNE | VHSPOOLGRIDSLSEQISFLEEQLGSCSCKKDS | /HSFQQLG | -510 -3777142 | GENERAL INFORMATION: | Ashkenazi, Avi Baker Kewin D | Botstein | Desnoyers, Eaton, Dan | Ferrara Filvaro | Fong, Sl | Gerber, | Goddard, Audrey | Godowsk | Gurney, | Kljavin | Kuo, Sophia (Napier, Marv | Pan, Jai | Faoni, Roy, Ma | Shelton Stewart | Tumas, Daniel Williams, P. | Wood, W | TITLE OF INVENTION: SECRETED AND IN TITLE OF INVENTION: Acids Encoding | FERENCE: P2630P1C18 APPLICATION NUMBER: US/09/978.681A | CURRENT FILING DATE: | APPLICATION NUMBER: 09/918585 FILING DATE: 2001-07-30 | CATION N | APPLICATION NUMBER: 60/064249 | FILLING UATE: 199/-1. APPLICATION NUMBER: | FILING DATE: 1997-11-1 APPLICATION NUMBER: 60 | FILING DATE: | G DATE: | APPLICATION N | APPLICATION N | CATION N | R FILING DATE: 1998-03-11 R APPLICATION NUMBER: 60/07 | IG DATE: | APPLICATION NUMBER: FILING DATE: 1998-0 | APPLICATION NUMBER: FILING DATE: 1998-0 | APPLICATION NUMBER: FILING DATE: 1998-0 |
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